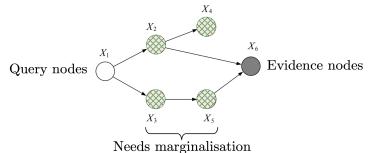
# CS5340 Uncertainty Modelling in AI

AY2022/23 Semester 2  $\cdot$  Prepared by Tian Xiao @snoidetx

CS5340 is about how to represent and reason with uncertainty in a computer.

## 5 Variable Elimination & Belief Propagation



0

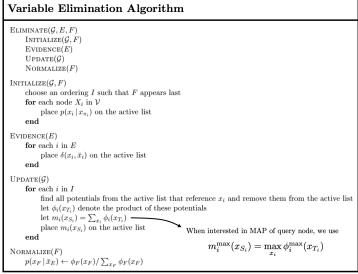
Variable Elimination: Use factorization and distributive law to reduce computational complexity:

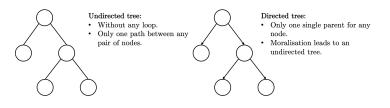
$$\begin{split} p(x_1, \overline{x}_6) \\ &= \sum_{x_2} \sum_{x_3} \sum_{x_4} \sum_{x_5} p(x_1) p(x_2 | x_1) p(x_3 | x_1) p(x_4 | x_2) p(x_5 | x_3) p(\overline{x}_6 | x_2, x_5) \\ &= p(x_1) \sum_{x_2} p(x_2 | x_1) \sum_{x_3} p(x_3 | x_1) \sum_{x_4} p(x_4 | x_2) \sum_{x_5} p(x_5 | x_3) p(\overline{x}_6 | x_2, x_5) \\ &= p(x_1) \sum_{x_2} p(x_2 | x_1) \sum_{x_3} p(x_3 | x_1) \sum_{x_4} p(x_4 | x_2) m_5(x_2, x_3) \\ &= p(x_1) \sum_{x_2} p(x_2 | x_1) \sum_{x_3} p(x_3 | x_1) m_5(x_2, x_3) \sum_{x_4} p(x_4 | x_2) \\ &= p(x_1) \sum_{x_2} p(x_2 | x_1) \sum_{x_3} p(x_3 | x_1) m_5(x_2, x_3) m_4(x_2) \\ &= p(x_1) \sum_{x_2} p(x_2 | x_1) m_4(x_2) \sum_{x_3} p(x_3 | x_1) m_5(x_2, x_3) \\ &= p(x_1) \sum_{x_2} p(x_2 | x_1) m_4(x_2) m_3(x_1, x_2) \\ &= p(x_1) m_2(x_1). \end{split}$$

- In this way,  $p(x_1|\overline{x}_6) = \frac{p(x_1)m_2(x_1)}{\sum\limits_{x_1} p(x_1)m_2(x_1)}$ .

- For UGMs, we do the same thing except changing local conditional probabilities into potentials of cliques.
  - ▶  $p(x_1|\overline{x}_6) = \frac{\frac{1}{Z}m_2(x_1)}{\frac{1}{Z}\sum_{x_1}m_2(x_1)}$ . The normalization factor Z in conditional probabilities, but **not** in marginal probabilities.
- Computational complexities: Analyzed through reconstituted graphs.
  - ▶ For UGMs, for each node X<sub>i</sub>, we connect all the remaining neighbours of X<sub>i</sub> and remove X<sub>i</sub>.
  - ▶ For DGMs, for each node X<sub>i</sub>, we connect all the parents of X<sub>i</sub>. In the end we drop the orientation of all edges (moralisation) and analyze it like UGMs.
  - ▶ Overall complexity:  $O(nk^M)$ , where M is the size of largest elimination clique.
  - ► Treewidth: One less than the smallest achievable cardinality of the largest elimination clique over all possible elimination orderings (NP-hard).
  - ▶ Heuristics:
    - \* Min-neighbors: Fewest number of dependent variables.

- \* Min-weight: Minimize product of cardinalities of variables.
- \* Min-fill: Minimize the size of the factor (elimination clique) that will be added.
- Limitations:
  - We have to re-run the variable elimination algorithm with every new query node.





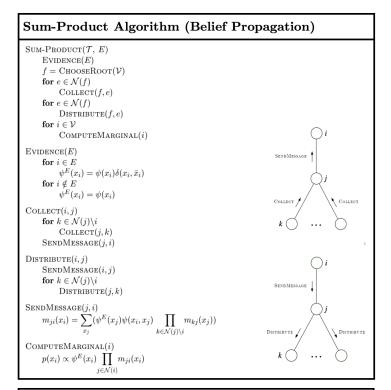
**Belief Propagation**: To obtain all marginals in the tree, we reuse messages to perform efficient inference.

- Works similarly for undirected and directed trees.
- Messages:

$$m_{ji}(x_i) = \sum_{x_j} \left( \psi^E(x_j) \psi(x_i, x_j) \prod_{k \in \mathcal{N}(j) \setminus i} m_{kj}(x_j) \right)$$
$$p(x_f | \overline{x}_E) \propto \psi^E(x_f) \prod_{e \in \mathcal{N}(f)} m_{ef}(x_f).$$

- Message-passing protocol: A node can send a message to a neighbouring node when and only when it has received messages from all of its other neighbours.
- MAP configurations: We record the maximising values in a table  $\delta_{ji}(x_i)$  when a message  $m_{ji}^{\max}(x_i)$  is sent from  $X_j$  to  $X_i$  (closer to root). We then use this table to define a consistent maximising configuration during an outward pass.
- Products of probabilities tend to underflow. We overcome this by using the monotone log scale:

$$\max_{x} p^{E}(x) = \max_{x} \log p^{E}(x).$$



### Max-Product Algorithm

MAX-PRODUCT $(\mathcal{T}, E)$ EVIDENCE(E) $f = CHOOSEROOT(\mathcal{V})$ for  $e \in \mathcal{N}(f)$ COLLECT(f, e) $MAP = \max_{x_f} (\psi^E(x_f) \prod_{e \in \mathcal{N}(f)} m_{ef}^{\max}(x_f))$  $x_f^* = \arg \max_{x_f} (\psi^E(x_f) \prod_{e \in \mathcal{N}(f)} m_{ef}^{\max}(x_f))$ for  $e \in \mathcal{N}(f)$ DISTRIBUTE(f, e)Collect(i, j)for  $k \in \mathcal{N}(j) \setminus i$ COLLECT(i, k)SENDMESSAGE(j, i)Distribute(i, j)SetValue(i, j)for  $k \in \mathcal{N}(j) \setminus i$ DISTRIBUTE(j, k)SENDMESSAGE(j, i) $m_{ji}^{\max}(x_i) = \max_{x_j}(\psi^E(x_j)\psi(x_i, x_j) \prod_{\substack{k \in \mathcal{N}(j) \setminus i \\ x_j \in \mathcal{N}(j) \setminus i}} m_{kj}^{\max}(x_j))$  $\delta_{ji}(x_i) \in \arg\max_{x_j}(\psi^E(x_j)\psi(x_i, x_j) \prod_{\substack{k \in \mathcal{N}(j) \setminus i \\ k \in \mathcal{N}(j) \setminus i}} m_{kj}^{\max}(x_j))$ SetValue(i, j) $x_j^* = \delta_{ji}(x_i^*)$ 



Polytree:Nodes with more than 1 parent.Moralisation leads to loops.

# 6 Factor Graph & Junction Tree

Factor Graph: Introduce additional nodes for the factors.

- Works for polytrees.
- A factor graph is a bipartite graph  $\mathcal{G}(\mathcal{V},\mathcal{F},\mathcal{E}),$  where
  - ▶  $\mathcal{V}$  is the set of random variables;
  - ►  $\mathcal{F}$  is the set of factors. In DGM, all the local conditional distributions  $p(x_i|x_{\pi_i})$  are represented as factors; in UGM, all the potential functions of cliques are represented as factors; normalising coefficients  $\frac{1}{Z}$  is a factor defined over the empty set of

variables;

- $\blacktriangleright \mathcal{E}$  is the set of all undirected edges.
- Two types of messages:
  - Messages  $\nu$  flow from variable to factor nodes:

$$\nu_{is}(x_i) = \prod_{t \in \mathcal{N}(i) \setminus s} \mu_{ti}(x_i)$$

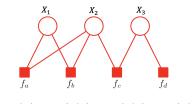
At leaf variable nodes,  $\nu_{is} = 1$ .

▶ Messages  $\mu$  flow from factor to variable nodes:

$$\mu_{si}(x_i) = \sum_{X_{\mathcal{N}(s)\setminus i}} \left( f_s\left(x_{\mathcal{N}(s)}\right) \prod_{j \in \mathcal{N}(s)\setminus i} \nu_{js}(x_j) \right)$$

At leaf factor nodes,  $\mu_{is} = f_s(x_i)$ .

- Message-passing protocol: A node can send a message to a neighbouring node when and only when it has received messages from all of its other neighbours (for both variable and factor nodes).
- $m_{ji}(x_i)$  in UGM is equal to  $\mu_{si}(x_i)$  in the factor graphs.



$$p(\mathbf{x}) = f_a(x_1, x_2) f_b(x_1, x_2) f_c(x_2, x_3) f_d(x_3)$$

Sum-Product Algorithm for Factor Graphs		
SUM-PRODUCT $(\mathcal{T}, E)$	$\mu$ -Distribute $(s, i)$	
EVIDENCE(E)	$\mu$ -SendMessage $(s, i)$	
$f = \text{CHOOSEROOT}(\mathcal{V})$	for $t \in \mathcal{N}(i) \backslash s$	
for $s \in \mathcal{N}(f)$	u-Distribute $(i, t)$	
$\mu$ -Collect $(f, s)$	$\nu$ -DISTRIBUTE $(i, s)$	
for $s \in \mathcal{N}(f)$	$\nu$ -SendMessAge $(i, s)$	
$\nu$ -DISTRIBUTE $(f, s)$	for $j \in \mathcal{N}(s) \setminus i$	
$\mathbf{for}i\in\mathcal{V}$	$\mu$ -DISTRIBUTE $(s, j)$	
COMPUTEMARGINAL(i)		
$\mu$ -Collect $(i, s)$	$\mu$ -SendMessAge $(s, i)$	
for $j \in \mathcal{N}(s) \setminus i$	$\mu_{si}(x_i) = \sum (f_s(x_{\mathcal{N}(s)}) \prod \nu_{js}(x_j))$	
$\nu$ -Collect $(s, j)$	$x_{\mathcal{N}(s)\setminus i}$ $j\in\overline{\mathcal{N}}(s)\setminus i$	
$\mu$ -SendMessage $(s, i)$	$\nu$ -SendMessage $(i, s)$	
	$ \nu_{is}(x_i) = \prod \mu_{ti}(x_i) $	
$\nu$ -Collect $(s, i)$	$t \in \mathcal{N}(i) \setminus s$	
for $t \in \mathcal{N}(i) \setminus s$	$C_{OMDUMP}M_{ADGINAL}(i)$	
$\mu$ -Collect $(i, t)$	COMPUTEMARGINAL(i)	
$\nu ext{-SendMessage}(i,s)$	$p(x_i) \propto  u_{is}(x_i) \mu_{si}(x_i)$	

**Junction Tree**: Probability distributions corresponding to loopy undirected graphs can be reparametrised as trees.

- Cluster graphs:
  - ▶ Nodes are clusters  $C_i \subseteq \{X_1, \dots, X_n\}$  where  $X_i$  are the random variables.
  - ▶ Edge between  $C_i$  and  $C_j$  associated with sepset  $S_{ij} = C_i \cap C_j$ .
- Family Preservation: Given a set of potentials  $\Psi \in \{\psi_1, \dots, \psi_k\}$ from an UGM, we assign each  $\psi_k$  to a cluster  $C_{\alpha(k)}$  such that  $\operatorname{Scope}[\psi_k] \subseteq C_{\alpha(k)}$ .
- Cluster potential:  $\phi_i(C_i) = \prod_{k:\alpha(k)=i} \psi_k.$
- Running Intersection Property: For each pair of clusters  $C_i, C_j$ and variables  $X \in C_i \cap C_j$ , there exists an unique path between  $C_i \cap C_j$  for which all clusters and sepsets contain X.

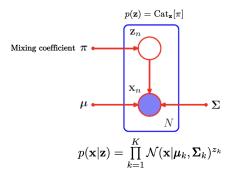
- ▶ Equivalently, for any X, the set of clusters and sepsets containing X from a tree.
- Cluster tree: A cluster graph without cycles is known as the a cluster tree.
- Clique trees (junction trees): A cluster tree that satisfies the running intersection property is called a clique tree or junction tree.
- Construction of junction trees:
  - 1. Triangulation via graph elimination.
  - Obtain clusters and all possible sepsets: use elimination cliques as clusters.
  - 3. Assign cluster potentials.
  - Get clique tree: find the maximum spanning tree with cardinality of sepsets as weight of edges. A cluster tree is a clique tree only if it is a maximal spanning tree.

#### 7 Mixture Models & Expectation Maximisation

**Gaussian Mixture Models (GMMs)**: GMMs can approximate almost any continuous density with arbitrary accuracy. It is a linear combination of Gaussian distributions:

$$p(\mathbf{x}) = \sum_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k),$$

- Each Gaussian density  $\mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$  is called a component of the mixture, and has its own mean  $(\boldsymbol{\mu}_k)$  and covariance  $(\boldsymbol{\Sigma}_k)$ .
- The parameters  $0 \le \pi_k \le 1$  is the mixing coefficients, and must sum to one:  $\sum_{k=1}^{K} \pi_k = 1$ .



- **Responsibility**: Measures the responsibility that component k takes for explaining the observation x:

$$\gamma(z_k) \equiv p(z_k = 1 | \mathbf{x}) = \frac{p(\mathbf{x} | z_k = 1)p(z_k = 1)}{\sum\limits_{j=1}^{K} p(\mathbf{x} | z_k = 1)p(z_k = 1)}$$
$$= \frac{\pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum\limits_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}.$$

**Expectation Maximisation for GMMs**: MLE for GMMs does not have a closed form solution, hence we derive an iterative solution.

- 1. Initialize the means  $\mu_k$ , covariances  $\Sigma_k$  and mixing coefficients  $\pi_k$ , and evaluate the initial value of the log likelihood.
- 2. Expectation Step: Evaluate the responsibilities  $\gamma(Z)$  using the current parameter values.

- 3. Maximization Step: Re-estimate the parameters using the current responsibilities.
- 4. Evaluate the log-likelihood and check for convergence.

**General EM Algorithm**: To find maximum likelihood solutions for models having latent variables:

$$\ln p(\mathbf{X}|\boldsymbol{\theta}) = \ln \int_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta}).$$

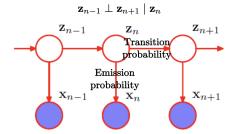
- 1. Choose an **initial setting** for the parameters  $\theta^{\text{old}}$ .
- 2. Expectation Step: Evaluate  $p(\mathbf{Z}|\mathbf{X}, \boldsymbol{\theta})$ .
- 3. Maximisation Step: Evaluate  $\boldsymbol{\theta}^{\text{new}}$  given by  $\boldsymbol{\theta}^{\text{new}} = \arg \max_{\boldsymbol{a}} \mathcal{Q}(\boldsymbol{\theta}, \boldsymbol{\theta}^{\text{old}})$ , where  $\mathcal{Q}(\boldsymbol{\theta}, \boldsymbol{\theta}^{\text{old}}) = \sum_{\mathbf{Z}} p(\mathbf{Z} | \mathbf{X}, \boldsymbol{\theta}^{\text{old}}) \ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\theta})$ .
- 4. Check for convergence of either the log likelihood or the parameter values, if not converged:  $\boldsymbol{\theta}^{\text{new}} \rightarrow \boldsymbol{\theta}^{\text{old}}$ .
- Theory behind EM:  $\ln p(\mathbf{X}|\boldsymbol{\theta}) = \mathcal{L}(q, \boldsymbol{\theta}) + \mathrm{KL}(q||p)$ , where

$$\blacktriangleright \mathcal{L}(q, \boldsymbol{\theta}) = \sum_{\mathbf{Z}} q(\mathbf{Z}) \ln \left\{ \frac{p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\theta})}{q(\mathbf{Z})} \right\};$$

• KL(q||p) = 
$$-\sum_{\mathbf{Z}} q(\mathbf{Z}) \ln \left\{ \frac{p(\mathbf{Z}|\mathbf{X}, \boldsymbol{\theta})}{q(\mathbf{Z})} \right\} \ge 0.$$

# 8 Hidden Markov Models

Latent variables form a Markov chain.

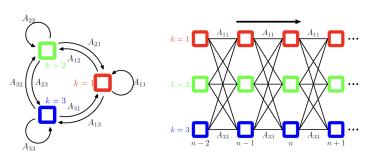


**Hidden Markov Models**: The Markov chain of latent variables gives rise to the graphical structure known as a state space model, where the joint distribution is given by

$$p(\mathbf{x}_1,\cdots,\mathbf{x}_n,\mathbf{z}_1,\cdots,\mathbf{z}_n) = p(\mathbf{z}_1) \left[\prod_{n=2}^N p(\mathbf{z}_n|\mathbf{z}_{n-1})\right] \prod_{n=1}^N p(\mathbf{x}_n|\mathbf{z}_n).$$

Transition diagram

Lattice representation



- Latent variables are discrete; observed variables can be either discrete or continuous.
- Transition probabilities: 1-of-K coding scheme for the discrete latent variables  $Z_n$ , which describes which mixture component is responsible for generating the observation  $X_n$ .
  - ▶  $p(z_n|z_{n-1})$  corresponds to a  $K \times K$  matrix **A** with the following properties:

1. 
$$A_{jk} = p(z_{nk} = 1 | z_{n-1,j} = 1).$$

- 2.  $0 \le A_{jk} \le 1$ , with  $\sum_{k} A_{jk} = 1$ .
- 3. K(K-1) independent parameters.
- $\blacktriangleright p(\mathbf{z}_n | \mathbf{z}_{n-1}, \mathbf{A}) = \prod_{k=1}^K \prod_{j=1}^K A_{jk}^{z_{n-1,j} z_{nk}}.$
- ► Initial latent variable  $Z_1$  does not have a parent node. It is represented as a categorical distribution  $p(\mathbf{z}_1 | \boldsymbol{\pi}) = \prod_{k=1}^{K} \pi_k^{z_{1k}}$ , where  $\sum_k \pi_k = 1$ .
- Emission probabilities:  $p(x_n|z_n, \phi)$ , where  $\phi$  is a set of parameters governing the distribution.
  - $p(\mathbf{x}_n | \mathbf{z}_n, \phi) = \prod_{k=1}^K p(\mathbf{x}_n | \phi_k)^{z_{nk}}.$
- Homogenous model: Only 1  ${\bf A}$  and  $\phi.$

#### Expectation Maximisation for HMMs:

- Marginal posterior distribution:  $\gamma(\mathbf{z}_n) = p(\mathbf{z}_n | \mathbf{X}, \boldsymbol{\theta}^{\text{old}}).$
- Joint posterior distribution:  $\xi(\mathbf{z}_{n-1}, \mathbf{z}_n) = p(\mathbf{z}_{n-1}, \mathbf{z}_n | \mathbf{X}, \boldsymbol{\theta}^{\text{old}})$ .
- 1. E Step: We use forward-backward algorithm:

# Forward-Backward Algorithm $\gamma(\mathbf{z}_n) = \frac{\alpha(\mathbf{z}_n)\beta(\mathbf{z}_n)}{p(\mathbf{X})}$ $\xi(\mathbf{z}_{n-1}, \mathbf{z}_n) = \frac{\alpha(\mathbf{z}_{n-1})p(\mathbf{x}_n|\mathbf{z}_n)p(\mathbf{z}_n|\mathbf{z}_{n-1})\beta(\mathbf{z}_n)}{p(\mathbf{X})}$ $\alpha(\mathbf{z}_n) = p(\mathbf{x}_n|\mathbf{z}_n)\sum_{\mathbf{z}_{n-1}}\alpha(\mathbf{z}_{n-1})p(\mathbf{z}_n|\mathbf{z}_{n-1})$ $\beta(\mathbf{z}_n) = \sum_{\mathbf{z}_{n+1}}\beta(\mathbf{z}_{n+1})p(\mathbf{x}_{n+1}|\mathbf{z}_{n+1})p(\mathbf{z}_{n+1}|\mathbf{z}_n)$

#### Forward-Backward Algorithm (Rescaled)

$$\begin{aligned} \gamma(\mathbf{z}_n) &= \widehat{\alpha}(\mathbf{z}_n)\widehat{\beta}(\mathbf{z}_n) \\ \xi(\mathbf{z}_{n-1}, \mathbf{z}_n) &= c_n \widehat{\alpha}(\mathbf{z}_{n-1}) p(\mathbf{x}_n | \mathbf{z}_n) p(\mathbf{z}_n | \mathbf{z}_{-1}) \widehat{\beta}(\mathbf{z}_n) \\ c_n \widehat{\alpha}(\mathbf{z}_n) &= p(\mathbf{x}_n | \mathbf{z}_n) \sum_{\mathbf{z}_{n-1}} \widehat{\alpha}(\mathbf{z}_{n-1}) p(\mathbf{z}_n | \mathbf{z}_{n-1}) \\ c_{n+1} \widehat{\beta}(\mathbf{z}_n) &= \sum_{\mathbf{z}_{n+1}} \widehat{\beta}(\mathbf{z}_{n+1}) p(\mathbf{x}_{n+1} | \mathbf{z}_{n+1}) p(\mathbf{z}_{n+1} | \mathbf{z}_n) \end{aligned}$$

2. M Step: Find  $\boldsymbol{\theta}$  that maximises

$$\mathcal{Q}(\boldsymbol{\theta}, \boldsymbol{\theta}^{\text{old}}) = \sum_{\mathbf{Z}} p(\mathbf{Z} | \mathbf{X}, \boldsymbol{\theta}^{\text{old}}) \ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\theta})$$
$$= \sum_{k=1}^{K} \gamma(z_{1k}) \ln \pi_k + \sum_{n=2}^{N} \sum_{j=1}^{K} \sum_{k=1}^{K} \xi(z_{n-1,j}, z_{nk}) \ln A_{jk} + \sum_{n=1}^{N} \sum_{k=1}^{K} \gamma(z_{nk}) \ln p(\mathbf{x}_n | \phi_k).$$

- Viterbi algorithm: Max-sum algorithm in factor trees.

#### 9 Monte-Carlo Inference

**Monte-Carlo Sampling**: Approximate a hard combinatorial problem by a much simpler problem using randomness.

- Monte-Carlo sampling is unbiased, consistent and converges at rate  $\frac{1}{\sqrt{N}}$ .
- Rejection sampling
- Importance sampling

Markov Chain Monte Carlo: Features adaptive proposals.

- Metropolis Hasting algorithm:

Algor	ithm : Metropolis-Hasting	
1. In	itialize $x^{(0)}$	
2. Fo	or $i = 0$ to $N - 1$	
3. 4.	Sample $u{\sim}\mathcal{U}_{[0,1]}$ Sample $x'{\sim}q(x' x^{(i)})$	// draw acceptance threshold // draw from proposal
5.	If $u < \mathcal{A}(x', x^{(i)}) = \min$	$\left\{1, rac{ ilde{p}(x')q(x^{(i)} x')}{ ilde{p}(x')q(x' x^{(i)})} ight\}$ // acceptance probability
6.	$x^{(i+1)} = x'$	// new sample is accepted
7. 8.	else $x^{(i+1)} = x^{(i)}$	// new sample is rejected // we create a duplicate of the previous sample

- Ergodic Theorem for Markov chains: If  $X_0, \dots, X_N$  is irreducible, homogenous, apeiodic discrete Markov chain with stationary distribution  $\pi$ , then

$$\frac{1}{N}\sum_{i=1}^{N}f(X_{i})\rightarrow \mathbb{E}[f(X)] \text{ as } N\rightarrow\infty$$

almost for sure where  $X \sim \pi$ , and

$$p(x_N = x | x_0) \to \pi(x) \ \forall x, x_0 \in \mathcal{X} \text{ as } N \to \infty.$$

- Stationary distribution:  $\pi T = \pi$ .
- Limiting distribution: Markov chain always converges to  $\pi$ .
- ► Irreducibility: For any state of the Markov chain, there is a positive probability of visiting all other states.
- ► Aperiodicity: The Markov chain should not get trapped in circles.
- Ergodicity: A Markov chain is ergodic if it is irreducible and aperiodic.
- ► Detailed balance: A probability vector  $\pi = p(x)$  satisfies detailed balance w.r.t. *T* if

$$\pi_a T_{ab} = \pi_b T_{ba}, \forall a, b \in \mathcal{X}.$$

Note that detailed balance implies stationary distribution.

**Gibbs Sampling**: A special case of the Metropolis Hasting algorithm where the acceptance probability is always one.

Algo	Algorithm : Gibbs Sampling		
	hitialize $\{x_i : i = 1,, M\}$ or $\tau = 1,, T$ :		
3. 4.	Sample $x_1^{\tau+1} \sim p(x_1   x_2^{(\tau)}, x_3^{(\tau)}, \dots, x_M^{(\tau)})$ . Sample $x_2^{\tau+1} \sim p(x_2   x_1^{(\tau+1)}, x_3^{(\tau)}, \dots, x_M^{(\tau)})$ .		
5.	Sample $x_j^{\tau+1} \sim p(x_j   x_1^{(\tau+1)},, x_{j-1}^{(\tau+1)}, x_{j+1}^{(\tau)},, x_M^{(\tau)}).$		
6.	Sample $x_M^{\tau+1} \sim p(x_M   x_1^{(\tau+1)}, x_2^{(\tau+1)}, \dots, x_{M-1}^{(\tau+1)})$		